# SEQUENCE LISTING

#### SENERAL INFORMATION:

- APPLICANT: HADASIT MEDICAL RESEARCH SERVICES AND DEVLOPMENT COMPANY (i)
- TITLE OF INVENTION: NOVEL HAPTOTACTIC PEPTIDES (ii)
- (iii) NUMBER OF SEQUENCES: 12

### CURRENT APPLICATION DATA:

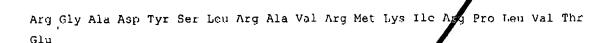
- (A) APPLICATION NUMBER: 09/487790
- (B) (B) FILING DATE: Jan. 20 2000
- (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE:
  - (2) INFORMATION FOR SEQ ID NO:1:
    - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 21
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Lys Gly Ser Trp Tyr Scr Met Arg Lys Met Ser Met Lys Ile Arg Pro Phe Pro Glu Glu

- (2) INFORMATION FOR SEQ 1D NO:2:
  - SEQUENCE CHARACTERISTICS: (i)
    - (A) LENGTH: 21
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Lys Thr Arg Trp Tyr∦Ser Met Lys Lys Thr Thr Met Lys Ile Ile Pro Phe Asn Arg Leu Thr

- (2) INFORMATION FOR SEQ ID NO:3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TO#OLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:



- (2) INFORMATION FOR SEQ TD NO:4:
  - SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Glu Ala Asp His Glu Gly Thr His Ser Thr Lys Arg Gly His Ala Lys Ser Arg Fro

- (2) INFORMATION FOR SEQ ID NO:5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ 1D NO:5:

Leu Thr Ile Gly Glu Gly Glu His His Leu Gly Gly Ala Lys Glu Ala Gly Asp Val

- (2) INFORMATION FOR SEQ ID NO:6:
  - (i) SEQUENCE CHARAGTERISTICS:
    - (A) LENGTH: 29
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRAPTION: SEQ ID NO:6:

X X Gly Val VaJ Trp X X X X Gly X X Tyr Ser X Arg X X Met Lys Ile Arg Fro X X X Glu

- (2) INFORMATION FOR SEQ ID NO:7:
  - SEQUENCE CHARACTERISTICS: (i)
    - (A) LENGTH: 65
    - (B) FYPE: nucleic acid
    - (C) STRANDEDNESS: double
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

AAGGGGTCATGGTATCAATGAGGAAGATGAGTATGAAGATCAGGCCCTTCTTCCCACAGCAATAG

- (2) INFORMATION FOR SEQ ID NO:8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (Λ) LENGTH: 63



- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

# AGAGGGGCAGATTATTCCCTCAGGGCTGTTCGCATGAAAATTAGGCCCCTTGTGACCCAATAG

(2) 'INFORMATION FOR SEQ ID NO:9:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 63
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: double
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

## AAAACCCGGTGGTATTCCATĞAAGAAAACCACTATGAAGATAATCCCATTCAACAGACTCACA

- (2) INFORMATION FOR SEQ ID NO:10
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS single
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ 1D NO:10:

Tyr Sor X Arg X X Met Lys le Arg Pro X X X Glu

# INFORMATION FOR SEQ ID NO: 11

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 20
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Lys X X Trp Tyr Ser Met X Lys X X Met Lys Ile X Pro Phe X X X

- (2) INFORMATION FOR SEQ ID NO: 12
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 30
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ 10 NO:12

Asp X Gly X X Trp X X Trp Lys X X Trp Tyr Ser Met X Lys X X Met Lys Ile X Pro Phe X X X X